DT12 Rec'd PCT/PT0 2 8 DEC 2004

Sequence Listing SEQUENCE LISTING

	<110>	Byrne, Barry J. Mah, Cathryn S.				
	<120> raav compositions and methods for delivery of human factor VII POLYPEPTIDES AND TREATMENT OF HEMOPHILIA A					
	<130>	4300.014300				
<140> UNKNOWN <141> 2004-12-28						
	<150> <151>	PCT/US03/20756 2003-06-30				
	<150> <151>	60/392,725 2002-06-28				
	<160>	17				
	<170>	PatentIn version 3.2				
	<210> <211> <212> <213>	1 1440 DNA Homo sapiens				
	<400> tcaacag	1 ggca ggggcagcac tgcagagatt tcatcatggt ctcccaggcc ctcaggctcc	60			
	tctgcc	ttct gcttgggctt cagggctgcc tggctgcagg cggggtcgct aaggcctcag	120			
	gaggaga	aaac acgggacatg ccgtggaagc cggggcctca cagagtcttc gtaacccagg	180			
	aggaag	ccca cggcgtcctg caccggcgcc ggcgcgccaa cgcgttcctg gaggagctgc	240			
	ggccgg	gctc cctggagagg gagtgcaagg aggagcagtg ctccttcgag gaggcccggg	300			
	agatct	tcaa ggacgcggag aggacgaagc tgttctggat ttcttacagt gatggggacc	360			
	agtgtg	cctc aagtccatgc cagaatgggg gctcctgcaa ggaccagctc cagtcctata	420			
	tctgct	tctg cctccctgcc ttcgagggcc ggaactgtga gacgcacaag gatgaccagc	480			
	tgatct	gtgt gaacgagaac ggcggctgtg agcagtactg cagtgaccac acgggcacca	540			
	agcgct	cctg tcggtgccac gaggggtact ctctgctggc agacggggtg tcctgcacac	600			
	ccacag	ttga atatccatgt ggaaaaatac ctattctaga aaaaagaaat gccagcaaac	660			
	cccaag	gccg aattgtgggg ggcaaggtgt gccccaaagg ggagtgtcca tggcaggtcc	720			
	tgttgt	tggt gaatggagct cagttgtgtg gggggaccct gatcaacacc atctgggtgg	780			
	tctccg	cggc ccactgtttc gacaaaatca agaactggag gaacctgatc gcggtgctgg	840			
	gcgagca	acga cctcagcgag cacgacgggg atgagcagag ccggcgggtg gcgcaggtca	900			
	tcatcc	ccag cacgtacgtc ccgggcacca ccaaccacga catcgcgctg ctccgcctgc	960			
	accagc	ccgt ggtcctcact gaccatgtgg tgcccctctg cctgcccgaa cggacgttct	1020			

ctgagaggac	gctggccttc	gtgcgcttct	cattggtcag	cggctggggc	cagctgctgg	1080
accgtggcgc	cacggccctg	gagctcatgg	tgctcaacgt	gccccggctg	atgacccagg	1140
actgcctgca	gcagtcacgg	aaggtgggag	actccccaaa	tatcacggag	tacatgttct	1200
gtgccggcta	ctcggatggc	agcaaggact	cctgcaaggg	ggacagtgga	ggcccacatg	1260
ccacccacta	ccggggcacg	tggtacctga	cgggcatcgt	cagctggggc	cagggctgcg	1320
caaccgtggg	ccactttggg	gtgtacacca	gggtctccca	gtacatcgag	tggctgcaaa	1380
agctcatgcg	ctcagagcca	cgcccaggag	tcctcctgcg	agccccattt	ccctagccca	1440

<210> 2

<211> 466

<212> PRT

<213> Homo sapiens

<400> 2

Met Val Ser Gln Ala Leu Arg Leu Leu Cys Leu Leu Leu Gly Leu Gln 10 15

Gly Cys Leu Ala Ala Gly Gly Val Ala Lys Ala Ser Gly Gly Glu Thr 20 25 30

Arg Asp Met Pro Trp Lys Pro Gly Pro His Arg Val Phe Val Thr Gln 35 40 45

Glu Glu Ala His Gly Val Leu His Arg Arg Arg Ala Asn Ala Phe $50 \hspace{1cm} 55 \hspace{1cm} 60$

Leu Glu Glu Leu Arg Pro Gly Ser Leu Glu Arg Glu Cys Lys Glu Glu 65 70 75 80

Gln Cys Ser Phe Glu Glu Ala Arg Glu Ile Phe Lys Asp Ala Glu Arg 85 90 95

Thr Lys Leu Phe Trp Ile Ser Tyr Ser Asp Gly Asp Gln Cys Ala Ser 100 105 110

Ser Pro Cys Gln Asn Gly Gly Ser Cys Lys Asp Gln Leu Gln Ser Tyr 115 120 125

Ile Cys Phe Cys Leu Pro Ala Phe Glu Gly Arg Asn Cys Glu Thr His 130 135 140

Lys Asp Asp Gln Leu Ile Cys Val Asn Glu Asn Gly Gly Cys Glu Gln 145 150 155 160

Tyr Cys Ser Asp His Thr Gly Thr Lys Arg Ser Cys Arg Cys His Glu Page 2 Gly Tyr Ser Leu Leu Ala Asp Gly Val Ser Cys Thr Pro Thr Val Glu 180 185 190 Tyr Pro Cys Gly Lys Ile Pro Ile Leu Glu Lys Arg Asn Ala Ser Lys 195 200 205 Pro Gln Gly Arg Ile Val Gly Gly Lys Val Cys Pro Lys Gly Glu Cys 210 220 Pro Trp Gln Val Leu Leu Leu Val Asn Gly Ala Gln Leu Cys Gly Gly 225 235 240 Thr Leu Ile Asn Thr Ile Trp Val Val Ser Ala Ala His Cys Phe Asp 245 250 255 Lys Ile Lys Asn Trp Arg Asn Leu Ile Ala Val Leu Gly Glu His Asp Leu Ser Glu His Asp Gly Asp Glu Gln Ser Arg Arg Val Ala Gln Val 275 280 285 Ile Ile Pro Ser Thr Tyr Val Pro Gly Thr Thr Asn His Asp Ile Ala 290 295 300 Leu Leu Arg Leu His Gln Pro Val Val Leu Thr Asp His Val Val Pro 305 310 Leu Cys Leu Pro Glu Arg Thr Phe Ser Glu Arg Thr Leu Ala Phe Val 325 330 335 Arg Phe Ser Leu Val Ser Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala 340 345 350 Thr Ala Leu Glu Leu Met Val Leu Asn Val Pro Arg Leu Met Thr Gln Asp Cys Leu Gln Gln Ser Arg Lys Val Gly Asp Ser Pro Asn Ile Thr 370 380 Glu Tyr Met Phe Cys Ala Gly Tyr Ser Asp Gly Ser Lys Asp Ser Cys 385 390 395 400 Lys Gly Asp Ser Gly Gly Pro His Ala Thr His Tyr Arg Gly Thr Trp $405 \hspace{1cm} 410 \hspace{1cm} 415$

Sequence Listing Tyr Leu Thr Gly Ile Val Ser Trp Gly Gln Gly Cys Ala Thr Val Gly 420 425 430

His Phe Gly Val Tyr Thr Arg Val Ser Gln Tyr Ile Glu Trp Leu Gln 435 440 445

Lys Leu Met Arg Ser Glu Pro Arg Pro Gly Val Leu Leu Arg Ala Pro 450 455 460

Phe Pro 465

<210> 3 <211> 1401 <212> DNA

<213> Homo sapiens

<400> 60 atggtctccc aggccctcag gctcctctgc cttctgcttg ggcttcaggg ctgcctggct 120 gcaggcgggg tcgctaaggc ctcaggagga gaaacacggg acatgccgtg gaagccgggg 180 cctcacagag tcttcgtaac ccaggaggaa gcccacggcg tcctgcaccg gcgccggcgc 240 gccaacgcgt tcctggagga gctgcggccg ggctccctgg agagggagtg caaggaggag 300 cagtgctcct tcgaggaggc ccgggagatc ttcaaggacg cggagaggac gaagctgttc 360 tggatttctt acagtgatgg ggaccagtgt gcctcaagtc catgccagaa tgggggctcc tgcaaggacc agctccagtc ctatatctgc ttctgcctcc ctgccttcga gggccggaac 420 tgtgagacgc acaaggatga ccagctgatc tgtgtgaacg agaacggcgg ctgtgagcag 480 tactgcagtg accacacggg caccaagcgc tcctgtcggt gccacgaggg gtactctctg 540 600 ctggcagacg gggtgtcctg cacacccaca gttgaatatc catgtggaaa aatacctatt ctagaaaaaa gaaatgccag caaaccccaa ggccgaattg tggggggcaa ggtgtgcccc 660 720 aaaggggagt gtccatggca ggtcctgttg ttggtgaatg gagctcagtt gtgtggggg 780 accctgatca acaccatctg ggtggtctcc gcggcccact gtttcgacaa aatcaagaac 840 tggaggaacc tgatcgcggt gctgggcgag cacgacctca gcgagcacga cggggatgag 900 cagagccggc gggtggcgca ggtcatcatc cccagcacgt acgtcccggg caccaccaac cacgacatcg cgctgctccg cctgcaccag cccgtggtcc tcactgacca tgtggtgccc 960 1020 ctctgcctgc ccgaacggac gttctctgag aggacgctgg ccttcgtgcg cttctcattg 1080 gtcagcggct ggggccagct gctggaccgt ggcgccacgg ccctggagct catggtgctc 1140 aacgtgcccc ggctgatgac ccaggactgc ctgcagcagt cacggaaggt gggagactcc ccaaatatca cggagtacat gttctgtgcc ggctactcgg atggcagcaa ggactcctgc 1200 1260 aagggggaca gtggaggccc acatgccacc cactaccggg gcacgtggta cctgacgggc Page 4

atcgtcagct	ggggccaggg	ctgcgcaacc	gtgggccact	ttggggtgta	caccagggtc	1320
tcccagtaca	tcgagtggct	gcaaaagctc	atgcgctcag	agccacgccc	aggagtcctc	1380
ctgcgagccc	catttcccta	g				1401

<210> 4

<211> 466

<212> PRT

<213> Homo sapiens

<400> 4

Met Val Ser Gln Ala Leu Arg Leu Leu Cys Leu Leu Gly Leu Gln 10 15

Gly Cys Leu Ala Ala Gly Gly Val Ala Lys Ala Ser Gly Gly Glu Thr 20 25 30

Arg Asp Met Pro Trp Lys Pro Gly Pro His Arg Val Phe Val Thr Gln 35 40 45

Glu Glu Ala His Gly Val Leu His Arg Arg Arg Ala Asn Ala Phe 50 60

Leu Glu Glu Leu Arg Pro Gly Ser Leu Glu Arg Glu Cys Lys Glu Glu 65 70 75 80

Gln Cys Ser Phe Glu Glu Ala Arg Glu Ile Phe Lys Asp Ala Glu Arg 85 90 95

Thr Lys Leu Phe Trp Ile Ser Tyr Ser Asp Gly Asp Gln Cys Ala Ser 100 105 110

Ser Pro Cys Gln Asn Gly Gly Ser Cys Lys Asp Gln Leu Gln Ser Tyr 115 120 125

Ile Cys Phe Cys Leu Pro Ala Phe Glu Gly Arg Asn Cys Glu Thr His 130 135 140

Lys Asp Asp Gln Leu Ile Cys Val Asn Glu Asn Gly Gly Cys Glu Gln 145 150 155 160

Tyr Cys Ser Asp His Thr Gly Thr Lys Arg Ser Cys Arg Cys His Glu 165 170 175

Gly Tyr Ser Leu Leu Ala Asp Gly Val Ser Cys Thr Pro Thr Val Glu 180 185 190

Sequence Listing Tyr Pro Cys Gly Lys Ile Pro Ile Leu Glu Lys Arg Asn Ala Ser Lys 195 200 205 Pro Gln Gly Arg Ile Val Gly Gly Lys Val Cys Pro Lys Gly Glu Cys 210 220 Pro Trp Gln Val Leu Leu Val Asn Gly Ala Gln Leu Cys Gly Gly 225 230 235 240 Thr Leu Ile Asn Thr Ile Trp Val Val Ser Ala Ala His Cys Phe Asp 245 250 255 Lys Ile Lys Asn Trp Arg Asn Leu Ile Ala Val Leu Gly Glu His Asp 260 265 270 Leu Ser Glu His Asp Gly Asp Glu Gln Ser Arg Arg Val Ala Gln Val 275 280 285 Ile Ile Pro Ser Thr Tyr Val Pro Gly Thr Thr Asn His Asp Ile Ala 290 295 300 Leu Leu Arg Leu His Gln Pro Val Val Leu Thr Asp His Val Val Pro 305 310 315 320Leu Cys Leu Pro Glu Arg Thr Phe Ser Glu Arg Thr Leu Ala Phe Val Arg Phe Ser Leu Val Ser Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala 340 350 Thr Ala Leu Glu Leu Met Val Leu Asn Val Pro Arg Leu Met Thr Gln Asp Cys Leu Gln Gln Ser Arg Lys Val Gly Asp Ser Pro Asn Ile Thr 370 375 380Glu Tyr Met Phe Cys Ala Gly Tyr Ser Asp Gly Ser Lys Asp Ser Cys 385 390 395 400 Lys Gly Asp Ser Gly Gly Pro His Ala Thr His Tyr Arg Gly Thr Trp 405 410 415 Tyr Leu Thr Gly Ile Val Ser Trp Gly Gln Gly Cys Ala Thr Val Gly 420 425 430 His Phe Gly Val Tyr Thr Arg Val Ser Gln Tyr Ile Glu Trp Leu Gln 435 440 445

Lys Leu Met Arg Ser Glu Pro Arg Pro Gly Val Leu Leu Arg Ala Pro 450 460

Phe Pro 465

<210> 5 <211> 1335 <212> DNA <213> Homo sapiens

<400> 5

atggtctccc	aggccctcag	gctcctctgc	cttctgcttg	ggcttcaggg	ctgcctggct	60
gcagtcttcg	taacccagga	ggaagcccac	ggcgtcctgc	accggcgccg	gcgcgccaac	120
gcgttcctgg	aggagctgcg	gccgggctcc	ctggagaggg	agtgcaagga	ggagcagtgc	180
tccttcgagg	aggcccggga	gatcttcaag	gacgcggaga	ggacgaagct	gttctggatt	240
tcttacagtg	atggggacca	gtgtgcctca	agtccatgcc	agaatggggg	ctcctgcaag	300
gaccagctcc	agtcctatat	ctgcttctgc	ctccctgcct	tcgagggccg	gaactgtgag	360
acgcacaagg	atgaccagct	gatctgtgtg	aacgagaacg	gcggctgtga	gcagtactgc	420
agtgaccaca	cgggcaccaa	gcgctcctgt	cggtgccacg	aggggtactc	tctgctggca	480
gacggggtgt	cctgcacacc	cacagttgaa	tatccatgtg	gaaaaatacc	tattctagaa	540
aaaagaaatg	ccagcaaacc	ccaaggccga	attgtggggg	gcaaggtgtg	ccccaaaggg	600
gagtgtccat	ggcaggtcct	gttgttggtg	aatggagctc	agttgtgtgg	ggggaccctg	660
atcaacacca	tctgggtggt	ctccgcggcc	cactgtttcg	acaaaatcaa	gaactggagg	720
aacctgatcg	cggtgctggg	cgagcacgac	ctcagcgagc	acgacgggga	tgagcagagc	780
cggcgggtgg	cgcaggtcat	catccccagc	acgtacgtcc	cgggcaccac	caaccacgac	840
atcgcgctgc	tccgcctgca	ccagcccgtg	gtcctcactg	accatgtggt	gcccctctgc	900
ctgcccgaac	ggacgttctc	tgagaggacg	ctggccttcg	tgcgcttctc	attggtcagc	960
ggctggggcc	agctgctgga	ccgtggcgcc	acggccctgg	agctcatggt	gctcaacgtg	1020
ccccggctga	tgacccagga	ctgcctgcag	cagtcacgga	aggtgggaga	ctccccaaat	1080
atcacggagt	acatgttctg	tgccggctac	tcggatggca	gcaaggactc	ctgcaagggg	1140
gacagtggag	gcccacatgc	cacccactac	cggggcacgt	ggtacctgac	gggcatcgtc	1200
agctggggcc	agggctgcgc	aaccgtgggc	cactttgggg	tgtacaccag	ggtctcccag	1260
tacatcgagt	ggctgcaaaa	gctcatgcgc	tcagagccac	gcccaggagt	cctcctgcga	1320
gccccatttc	cctag					1335

<211> 444

<212> PRT

<213> Homo sapiens

<400> 6

Met Val Ser Gln Ala Leu Arg Leu Leu Cys Leu Leu Gly Leu Gln 1 5 10 15

Gly Cys Leu Ala Ala Val Phe Val Thr Gln Glu Glu Ala His Gly Val 20 25 30

Leu His Arg Arg Arg Ala Asn Ala Phe Leu Glu Glu Leu Arg Pro 35 40 45

Gly Ser Leu Glu Arg Glu Cys Lys Glu Glu Gln Cys Ser Phe Glu Glu 50 60

Ala Arg Glu Ile Phe Lys Asp Ala Glu Arg Thr Lys Leu Phe Trp Ile 65 70 75 80

Ser Tyr Ser Asp Gly Asp Gln Cys Ala Ser Ser Pro Cys Gln Asn Gly 85 90 95

Gly Ser Cys Lys Asp Gln Leu Gln Ser Tyr Ile Cys Phe Cys Leu Pro 100 105 110

Ala Phe Glu Gly Arg Asn Cys Glu Thr His Lys Asp Asp Gln Leu Ile 115 120 125

Cys Val Asn Glu Asn Gly Gly Cys Glu Gln Tyr Cys Ser Asp His Thr 130 135 140

Gly Thr Lys Arg Ser Cys Arg Cys His Glu Gly Tyr Ser Leu Leu Ala 145 150 155 160

Asp Gly Val Ser Cys Thr Pro Thr Val Glu Tyr Pro Cys Gly Lys Ile 165 170 175

Pro Ile Leu Glu Lys Arg Asn Ala Ser Lys Pro Gln Gly Arg Ile Val 180 185 190

Gly Gly Lys Val Cys Pro Lys Gly Glu Cys Pro Trp Gln Val Leu Leu 195 200 205

Leu Val Asn Gly Ala Gln Leu Cys Gly Gly Thr Leu Ile Asn Thr Ile 210 215 220

Trp Val Val Ser Ala Ala His Cys Phe Asp Lys Ile Lys Asn Trp Arg Page 8 Asn Leu Ile Ala Val Leu Gly Glu His Asp Leu Ser Glu His Asp Gly 245 250 255

Asp Glu Gln Ser Arg Arg Val Ala Gln Val Ile Ile Pro Ser Thr Tyr 260 265 270

Val Pro Gly Thr Thr Asn His Asp Ile Ala Leu Leu Arg Leu His Gln 275 280 285

Pro Val Val Leu Thr Asp His Val Val Pro Leu Cys Leu Pro Glu Arg 290 295 300

Thr Phe Ser Glu Arg Thr Leu Ala Phe Val Arg Phe Ser Leu Val Ser 305 310 315 320

Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala Thr Ala Leu Glu Leu Met 325 330 335

Val Leu Asn Val Pro Arg Leu Met Thr Gln Asp Cys Leu Gln Gln Ser 340 345 350

Arg Lys Val Gly Asp Ser Pro Asn Ile Thr Glu Tyr Met Phe Cys Ala 355 360 365

Gly Tyr Ser Asp Gly Ser Lys Asp Ser Cys Lys Gly Asp Ser Gly Gly 370 380

Pro His Ala Thr His Tyr Arg Gly Thr Trp Tyr Leu Thr Gly Ile Val 385 390 395 400

Ser Trp Gly Gln Gly Cys Ala Thr Val Gly His Phe Gly Val Tyr Thr 405 410 415

Arg Val Ser Gln Tyr Ile Glu Trp Leu Gln Lys Leu Met Arg Ser Glu 420 425 430

Pro Arg Pro Gly Val Leu Leu Arg Ala Pro Phe Pro 435 440

<210> 7

<211> 1341

<212> DNA

<213> Rattus norvegicus

<400> 7

atggttccac agactcacgg actgctgctt ctctactttc tgctccagct ccagggaccc

Sequence Listing ctaggggctg tggttttcat aacccaggag gaagcacacg gtgtcctaca caggcaaagg 120 180 cgtgccaact cactcctaga ggagctttgg tctagctcct tggagaggga gtgcaatgaa 240 gagcggtgct cctttgagga ggcccgagag atcttcaaga gccctgagag aaccaagcag ttctggacta tttacagcga tggcgaccag tgtgcctcga atccatgtca gaacgggggt 300 360 acctgccagg atcacctcaa gtcttatgtc tgcttctgcc ccctagactt tgagggccgg 420 aactgtgaga aaaacaagaa tgagcagctg atctgtgcaa atgaaaatgg tgactgtgac 480 cagtactgca gggaccacgt agggaccaag cgtacctgta gctgtcacga ggactacgtg 540 ctgcagccag atgaggtgtc ctgcaaacca aaagttgagt acccatgcgg gagaatacct 600 gttgtagaaa aaagaaactt cagcagaccc caaggccgga ttgtgggagg ctatgtgtgc cccaaagggg agtgcccatg gcaggctgtg ctgaaattca atgaggcatt gctgtgtggg 660 gccgtcctgc tggacaccag atggatagta actgcagccc actgcttcga taaattcggg 720 780 aaattggtaa acatcacagt ggtgttgggt gaacacgact tcagtgagaa ggaggggact 840 gagcaagtac ggctggtgga acaggtcatc atgcccaaca agtacacccg cggcaggact 900 gaccatgaca tcgccctggt ccgccttcac cggcctgtaa ccttcactga ctacgtggta 960 cctctgtgtc tgcctgaacg ggccttctcc gagaacaccc tagccagcat ccgcttctcg 1020 agggtcagcg gctggggcca gctactggac cgtggtgcca cagctctgga gctcatggtc 1080 atcgaggtgc cccggctgat gacccaggac tgcctggagc atgccaaaca cagtgctaac 1140 acccccagaa tcacggagaa catgttctgc gccggctaca tggacggcac caaggacgcc 1200 tgcaagggtg acagtggagg cccacacgcc acccactacc atggcacttg gtatctgaca 1260 ggtgtggtca gctgggggga gggctgtgca gctatcggcc acatcggggt gtacaccagg 1320 gtctcccagt acatagactg gctggtcaaa tacatggact ccaagctccg ggttgggatt

<210> 8 <211> 446

<213> Rattus norvegicus

tctcgagtct ccctactgta g

<400> 8

Met Val Pro Gln Thr His Gly Leu Leu Leu Leu Tyr Phe Leu Leu Gln 10 15

Leu Gln Gly Pro Leu Gly Ala Val Val Phe Ile Thr Gln Glu Glu Ala 20 25 30

His Gly Val Leu His Arg Gln Arg Arg Ala Asn Ser Leu Leu Glu Glu 35 40 45

1341

Leu Trp Ser Ser Ser Leu Glu Arg Glu Cys Asn Glu Glu Arg Cys Ser 50 60 Phe Glu Glu Ala Arg Glu Ile Phe Lys Ser Pro Glu Arg Thr Lys Gln 65 70 75 80 Phe Trp Thr Ile Tyr Ser Asp Gly Asp Gln Cys Ala Ser Asn Pro Cys 85 90 95 Gln Asn Gly Gly Thr Cys Gln Asp His Leu Lys Ser Tyr Val Cys Phe 100 105 110 Cys Pro Leu Asp Phe Glu Gly Arg Asn Cys Glu Lys Asn Lys Asn Glu 115 120 125 Gln Leu Ile Cys Ala Asn Glu Asn Gly Asp Cys Asp Gln Tyr Cys Arg 130 135 140 Asp His Val Gly Thr Lys Arg Thr Cys Ser Cys His Glu Asp Tyr Val 145 150 155 160 Leu Gln Pro Asp Glu Val Ser Cys Lys Pro Lys Val Glu Tyr Pro Cys 165 170 175 Gly Arg Ile Pro Val Val Glu Lys Arg Asn Phe Ser Arg Pro Gln Gly 180 185 190 Arg Ile Val Gly Gly Tyr Val Cys Pro Lys Gly Glu Cys Pro Trp Gln
195 200 205 Ala Val Leu Lys Phe Asn Glu Ala Leu Leu Cys Gly Ala Val Leu Leu 210 215 220 Asp Thr Arg Trp Ile Val Thr Ala Ala His Cys Phe Asp Lys Phe Gly 235 235 240 Lys Leu Val Asn Ile Thr Val Val Leu Gly Glu His Asp Phe Ser Glu 245 250 255 Lys Glu Gly Thr Glu Gln Val Arg Leu Val Glu Gln Val Ile Met Pro 260 265 270 Asn Lys Tyr Thr Arg Gly Arg Thr Asp His Asp Ile Ala Leu Val Arg 275 280 285 Leu His Arg Pro Val Thr Phe Thr Asp Tyr Val Val Pro Leu Cys Leu 290 295 300 Page 11

Pro Glu Arg Ala Phe Ser Glu Asn Thr Leu Ala Ser Ile Arg Phe Ser 305 310 315 320
Arg Val Ser Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala Thr Ala Leu 325 330 335
Glu Leu Met Val Ile Glu Val Pro Arg Leu Met Thr Gln Asp Cys Leu 340 345 350
Glu His Ala Lys His Ser Ala Asn Thr Pro Arg Ile Thr Glu Asn Met 355 360 365
Phe Cys Ala Gly Tyr Met Asp Gly Thr Lys Asp Ala Cys Lys Gly Asp 370 380
Ser Gly Gly Pro His Ala Thr His Tyr His Gly Thr Trp Tyr Leu Thr 385 390 395 400
Gly Val Val Ser Trp Gly Glu Gly Cys Ala Ala Ile Gly His Ile Gly 405 410 415
Val Tyr Thr Arg Val Ser Gln Tyr Ile Asp Trp Leu Val Lys Tyr Met 420 425 430
Asp Ser Lys Leu Arg Val Gly Ile Ser Arg Val Ser Leu Leu 435 440 445
<210> 9 <211> 1671 <212> DNA <213> Danio rerio
<400> 9 atgagtctgc tgcttgtgtt ttctctgctc tggagtctcc attactgcca ttcagcagca 6
gtgttcgtgc acagagatga agctcacgag gtgttgatca ggagcaaaag agccaactca 12
ggctggtttg aggagctgaa gacggggaat ctggagcgcg agtgtctgga ggagaaatgc 18
tcgtatgagg aggcgcgcga ggtgttcgag cacacagagg ccacgaatga gttctggaag 24
atctacgatg ttaaagatca ctgcgcatcc agtccatgtg agcatgacgg gctctgcacc 30
acacagaacg cggactccta catgtgtttg tgtgcgccgg gcttcagcgg acgccactgt 36
gagcaatcga ttggagacgt tctcgactcc tgtctgcatg ataacggcgg ctgcgaacac 420
ttctgcacgg agcaggacgg acggagaaac tgctcctgcg cagacgggta ttacctagat 48
aacagcgggc agaagtgccg gagtcacgag gtgtttccat gtgggaaggt tcctctcctg 540
caggctggaa aagctgcgga tcatcaggtg gatctcagat ctcgtatcgt tggaggatct 600 Page 12

gaatgtccta	aaggtcactg	tccgtggcag	gtgctgctga	agtacggtga	gaagggtttc	660
tgtggaggtg	tgatctacaa	gcccacctgg	atcctcacag	ctgctcactg	cttggaaaag	720
ctcaaggtca	agttcctcag	gatagtggca	ggtgagcatg	atctggaggt	ggacgagggc	780
acggagcagc	tcatccaggt	ggatcagatg	ttcacacacc	ctgcgtacgt	gtctgagaca	840
gcggacagtg	acatcgccct	gctgcgtctg	cgcaccccca	tcgtctacag	tgtgtatgcg	900
gtgccggtgt	gtttgccgct	gcgggagatg	gcggagcgcg	agctgtgggc	ggtcagcaaa	960
cacacggtga	gcggctgggg	caaacgcagc	gaggacgggc	cgacctctcg	cctgctgcgc	1020
cggctgctgg	tgccgcgcat	ccgcacgcag	gagtgtgtgc	aggtcagcaa	cctcacgctc	1080
accagcaaca	tgttctgcgc	cggatacatc	gagggccggc	aggactcctg	taagggtgac	1140
agcggcggcc	cgctggtgac	ccggtaccga	gacaccgcct	tcctactggg	catcgtgagc	1200
tgggggaaag	gctgcgctcg	cccgggctcc	tacggcatct	acacacgcgt	gtccaactac	1260
ctgcagtgga	tccgacaaac	aaccaacacc	acgatacact	gatgaagaca	tgacccgggt	1320
gcattgctca	tcaagattgc	tactcttagg	tgaacaatta	acaaatatta	actattatag	1380
ttaatgtttg	taaaaaatag	caaaattata	ttgaaaataa	aaaatattta	tattaattat	1440
gaagtgacgg	cgattacttt	aattatccaa	gacggtgtta	tagcccaaaa	tacccaatag	1500
ttgagcatca	gctgctttcc	tgacatcctg	tacatattag	actcggatct	gatattttgc	1560
acaggttata	ttgcattttt	agcaggtatt	taatgatttt	gctctgatta	atcaggagat	1620
gtgcagctca	ttatctccat	attattaatg	ctcaactgta	gtaaacactc	g	1671

<210>

433

<212> PRT

<213> Danio rerio

<400> 10

Met Ser Leu Leu Leu Val Phe Ser Leu Leu Trp Ser Leu His Tyr Cys $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

His Ser Ala Ala Val Phe Val His Arg Asp Glu Ala His Glu Val Leu 20 25 30

Ile Arg Ser Lys Arg Ala Asn Ser Gly Trp Phe Glu Glu Leu Lys Thr 35 40 45

Gly Asn Leu Glu Arg Glu Cys Leu Glu Glu Lys Cys Ser Tyr Glu Glu 50 60

Ala Arg Glu Val Phe Glu His Thr Glu Ala Thr Asn Glu Phe Trp Lys 70 75 80 Page 13

Ile Tyr Asp Val Lys Asp His Cys Ala Ser Ser Pro Cys Glu His Asp 85 90 95 Gly Leu Cys Thr Thr Gln Asn Ala Asp Ser Tyr Met Cys Leu Cys Ala 100 105 110 Pro Gly Phe Ser Gly Arg His Cys Glu Gln Ser Ile Gly Asp Val Leu 115 120 125 Asp Ser Cys Leu His Asp Asn Gly Gly Cys Glu His Phe Cys Thr Glu 130 135 140 Gln Asp Gly Arg Arg Asn Cys Ser Cys Ala Asp Gly Tyr Tyr Leu Asp 145 150 155 160 Asn Ser Gly Gln Lys Cys Arg Ser His Glu Val Phe Pro Cys Gly Lys 165 170 175 Val Pro Leu Leu Gln Ala Gly Lys Ala Ala Asp His Gln Val Asp Leu 180 185 190 Arg Ser Arg Ile Val Gly Gly Ser Glu Cys Pro Lys Gly His Cys Pro 195 200 205 Gln Val Leu Leu Lys Tyr Gly Glu Lys Gly Phe Cys Gly Gly Val 210 225 220 Ile Tyr Lys Pro Thr Trp Ile Leu Thr Ala Ala His Cys Leu Glu Lys 235 230 240 Leu Lys Val Lys Phe Leu Arg Ile Val Ala Gly Glu His Asp Leu Glu 245 250 255 Val Asp Glu Gly Thr Glu Gln Leu Ile Gln Val Asp Gln Met Phe Thr 260 265 270 His Pro Ala Tyr Val Ser Glu Thr Ala Asp Ser Asp Ile Ala Leu Leu 275 280 285 Arg Leu Arg Thr Pro Ile Val Tyr Ser Val Tyr Ala Val Pro Val Cys Leu Pro Leu Arg Glu Met Ala Glu Arg Glu Leu Trp Ala Val Ser Lys 305 310 His Thr Val Ser Gly Trp Gly Lys Arg Ser Glu Asp Gly Pro Thr Ser Page 14

Arg Leu Leu Arg Arg Leu Leu Val Pro Arg Ile Arg Thr Gln Glu Cys 340 345 350

Val Gln Val Ser Asn Leu Thr Leu Thr Ser Asn Met Phe Cys Ala Gly 355 360 365

Tyr Ile Glu Gly Arg Gln Asp Ser Cys Lys Gly Asp Ser Gly Gly Pro 370 380

Leu Val Thr Arg Tyr Arg Asp Thr Ala Phe Leu Leu Gly Ile Val Ser 385 390 395 400

Trp Gly Lys Gly Cys Ala Arg Pro Gly Ser Tyr Gly Ile Tyr Thr Arg 405 410 415

Val Ser Asn Tyr Leu Gln Trp Ile Arg Gln Thr Thr Asn Thr Thr Ile 420 425 430

His

<210> 11

<211> 1341 <212> DNA

<213> Mus musculus

<400> 11

60 atggttccac aggcgcatgg gctgctgctt ctctgctttc tgctccagct ccagggacct ctagggactg cagttttcat aacccaggag gaagcacatg gtgtcctaca caggcaaagg 120 180 cgtgccaact cactcctgga ggagctttgg cccggctctc tggagagaga gtgcaatgag 240 gaacagtgct cctttgagga ggcccgggag atcttcaaga gccctgagag gaccaagcag ttctggattg tttacagtga tggggaccag tgtgcctcga atccatgtca gaacgtaggt 300 360 acctgccagg atcatctcaa gtcttacgtc tgcttctgcc tcctagactt tgagggtcgg 420 aactgtgaga aaagcaagaa tgagcagctg atctgtgcaa atgaaaatgg tgactgtgac 480 cagtactgca gggaccatgt agggaccaag cgtacctgta gctgtcatga ggactacacg 540 ctacagccag atgaggtgtc ctgcaaacca aaagttgagt acccgtgtgg gagaatacct gttgtagaaa aaagaaactc cagcagccgc caaggccgca ttgtgggagg caacgtgtgc 600 660 cccaaagggg agtgtccatg gcaggctgtg ctgaaaatca atgggttatt gctgtgtggg gccgtcctgc tggacgccag atggatagtg accgcagccc actgcttcga taatatccgc 720 780 tactggggaa acatcacagt ggtgatgggt gaacatgact tcagtgagaa ggatggggat

aaacaaatac	aa.caaataa.c	26266555	Sequence L		69963393+6	940
gagcaagtac	gacgggtgac	acayyccacc	atycccyaca	agracaticty	cyycaayacc	840
aaccacgaca	ttgccctgct	ccgccttcac	cggcctgtga	ccttcactga	ctacgtggtg	900
cccctgtgtc	tgcctgaaaa	gtccttctcc	gagaacaccc	tagccagaat	ccgcttctca	960
agggtcagtg	gctggggcca	gctactggac	cgtggtgcca	cagccctgga	actcatgtcc	1020
atcgaggtgc	cccggctgat	gacccaggac	tgtctggagc	acgccaagca	cagctctaac	1080
acccccaaaa	tcacagagaa	catgttctgc	gctggctaca	tggatggtac	caaggacgcc	1140
tgcaagggtg	acagcggtgg	cccacatgcc	acgcactacc	atggcacatg	gtatctgaca	1200
ggtgtggtca	gctgggggga	gggctgtgca	gctattggtc	acattggggt	gtacaccagg	1260
gtctcccagt	acatagactg	gctggtcaga	cacatggact	ccaagctcca	ggttggggtt	1320
ttccgactcc	cactactgta	g				1341

<210> 12

<211> 446 <212> PRT

<213> Mus musculus

<400> 12

Met Val Pro Gln Ala His Gly Leu Leu Leu Leu Cys Phe Leu Leu Gln 10 15

Leu Gln Gly Pro Leu Gly Thr Ala Val Phe Ile Thr Gln Glu Glu Ala 20 25 30

His Gly Val Leu His Arg Gln Arg Arg Ala Asn Ser Leu Leu Glu Glu 35 40 45

Leu Trp Pro Gly Ser Leu Glu Arg Glu Cys Asn Glu Glu Gln Cys Ser 50 60

Phe Glu Glu Ala Arg Glu Ile Phe Lys Ser Pro Glu Arg Thr Lys Gln 65 70 75 80

Phe Trp Ile Val Tyr Ser Asp Gly Asp Gln Cys Ala Ser Asn Pro Cys 85 90 95

Gln Asn Val Gly Thr Cys Gln Asp His Leu Lys Ser Tyr Val Cys Phe $100 \hspace{1cm} 105 \hspace{1cm} 110$

Cys Leu Leu Asp Phe Glu Gly Arg Asn Cys Glu Lys Ser Lys Asn Glu 115 120 125

Gln Leu Ile Cys Ala Asn Glu Asn Gly Asp Cys Asp Gln Tyr Cys Arg 130 135 140

Asp His Val Gly Thr Lys Arg Thr Cys Ser Cys His Glu Asp Tyr Thr 145 150 155 160 Leu Gln Pro Asp Glu Val Ser Cys Lys Pro Lys Val Glu Tyr Pro Cys 165 170 175 Gly Arg Ile Pro Val Val Glu Lys Arg Asn Ser Ser Ser Arg Gln Gly 180 185 190 Arg Ile Val Gly Gly Asn Val Cys Pro Lys Gly Glu Cys Pro Trp Gln
195 200 205 Ala Val Leu Lys Ile Asn Gly Leu Leu Leu Cys Gly Ala Val Leu Leu 210 220 Asp Ala Arg Trp Ile Val Thr Ala Ala His Cys Phe Asp Asn Ile Arg 225 230 235 240 Tyr Trp Gly Asn Ile Thr Val Val Met Gly Glu His Asp Phe Ser Glu 245 250 255 Lys Asp Gly Asp Glu Gln Val Arg Arg Val Thr Gln Val Ile Met Pro 260 265 270 Asp Lys Tyr Ile Arg Gly Lys Ile Asn His Asp Ile Ala Leu Leu Arg 275 280 285 Leu His Arg Pro Val Thr Phe Thr Asp Tyr Val Val Pro Leu Cys Leu Pro Glu Lys Ser Phe Ser Glu Asn Thr Leu Ala Arg Ile Arg Phe Ser 305 310 315 Arg Val Ser Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala Thr Ala Leu 325 330 335 Glu Leu Met Ser Ile Glu Val Pro Arg Leu Met Thr Gln Asp Cys Leu 340 350Glu His Ala Lys His Ser Ser Asn Thr Pro Lys Ile Thr Glu Asn Met $355 \hspace{1.5cm} 360 \hspace{1.5cm} 365$ Phe Cys Ala Gly Tyr Met Asp Gly Thr Lys Asp Ala Cys Lys Gly Asp 370 375 380 Ser Gly Gly Pro His Ala Thr His Tyr His Gly Thr Trp Tyr Leu Thr 385 390 395 400 Page 17

Gly Val Val Ser Trp Gly Glu Gly Cys Ala Ala Ile Gly His Ile Gly 405 410 415

Val Tyr Thr Arg Val Ser Gln Tyr Ile Asp Trp Leu Val Arg His Met 420 425 430

Asp Ser Lys Leu Gln Val Gly Val Phe Arg Leu Pro Leu Leu 435 440 445

<210> 13 <211> 1260

<212> DNA

<213> Gallus gallus

<400> 13

60 atgqtttcca ggcagtgcgt ggctttgctg ctctgcttcc cgctgctggt tcctccttct 120 ctggaagcag tctttttaaa gcaggaagag gcaaacagca tttttcaaag gcacagaaga gccaatagct tctttgaaga gataaagctg gggccactag agcgagaatg catagaagaa 180 aagtgttcat ttgaggaagc aagagagatc taccgtgatg atgagaggac aaaagagttc 240 tggcacatct attctgaccc caaccagtgt gactccagcc cctgtcagaa cggagggagc 300 360 tgcgatgacc agtttcagga ttatgtctgc cgctgtcctc cggagtacga gggcaaaagc 420 tgtgaaacag ctgtggccga gaacctgaag tgcatttacg acaacggcgg ctgtgagcag 480 tactgtgctg acgagcagtc tgaaaaacga gtgtgcttct gtgcagaggg ctacgcttta 540 gcgagtgatg gagtgtcctg cattccccaa gtgaaatacc cttgtggaac gataccagtg 600 ctggcaagaa agaatacaac tgctcagggg agaatagtag gtggtgtcac ctgtcctccg 660 ggtgaatgtc catggcaagc ccttataata caggatcaga aagggaaatg tgggggtagt 720 ctgctctcac cagagtgggt ggtgactgca gctcattgcc tggactacgc tcattccaaa cagctccggg tgaggctggg tgaatactca gtaaaagttg ctgagaaaac tgagcaagaa 780 840 agtggagtta gcaagatcat caggcacgaa gaatacacca ttggacaagt caatcatgac 900 attgccctcc tgaagctgga aacacccgtg aatctcaccg atttcgttgt gccaatatgt ttgcctgaaa aacggtttgc agtgtacgag ctgtcctcca ttaagttctc aatggtgagc 960 1020 ggatggggac ggctactaga tggaggggct acttctactt ttctgatgcg agttcatttg 1080 ccccgtgtaa agacacaaga atgtgaaaag caggctaatt tgaacatcac cgagaatatg 1140 ttctgtgcag gagacctgac cggtaaaaaa gactcctgca agggagacag tggtggacct 1200 cacgctacaa agtacaagaa cacctggttt ctgactggga ttgtcagctg gggaaagggt tgtgctgttg aaggcagcta cggggtgtac acaagggtat ccaqatacat caactggttg 1260

<210> 14

<211> 425

<212> PRT <213> Gallus gallus

<400> 14

Met Val Ser Arg Gln Cys Val Ala Leu Leu Cys Phe Pro Leu Leu 1 5 10 15

Val Pro Pro Ser Leu Glu Ala Val Phe Leu Lys Gln Glu Glu Ala Asn 20 25 30

Ser Ile Phe Gln Arg His Arg Arg Ala Asn Ser Phe Phe Glu Glu Ile 35 40 45

Lys Leu Gly Pro Leu Glu Arg Glu Cys Ile Glu Glu Lys Cys Ser Phe 50 55 60

Glu Glu Ala Arg Glu Ile Tyr Arg Asp Asp Glu Arg Thr Lys Glu Phe 65 70 75 80

Trp His Ile Tyr Ser Asp Pro Asn Gln Cys Asp Ser Ser Pro Cys Gln 85 90 95

Asn Gly Gly Ser Cys Asp Asp Gln Phe Gln Asp Tyr Val Cys Arg Cys 100 105 110

Pro Pro Glu Tyr Glu Gly Lys Ser Cys Glu Thr Ala Val Ala Glu Asn 115 120 125

Leu Lys Cys Ile Tyr Asp Asn Gly Gly Cys Glu Gln Tyr Cys Ala Asp 130 140

Glu Gln Ser Glu Lys Arg Val Cys Phe Cys Ala Glu Gly Tyr Ala Leu 145 150 155 160

Ala Ser Asp Gly Val Ser Cys Ile Pro Gln Val Lys Tyr Pro Cys Gly 165 170 175

Thr Ile Pro Val Leu Ala Arg Lys Asn Thr Thr Ala Gln Gly Arg Ile 180 185 190

Val Gly Gly Val Thr Cys Pro Pro Gly Glu Cys Pro Trp Gln Ala Leu 195 200 205

Ile Ile Gln Asp Gln Lys Gly Lys Cys Gly Gly Ser Leu Leu Ser Pro
210 215 220

Sequence Listing Glu Trp Val Val Thr Ala Ala His Cys Leu Asp Tyr Ala His Ser Lys 225 230 235 240 Gln Leu Arg Val Arg Leu Gly Glu Tyr Ser Val Lys Val Ala Glu Lys 245 250 255 Thr Glu Gln Glu Ser Gly Val Ser Lys Ile Ile Arg His Glu Glu Tyr 260 265 270 Thr Ile Gly Gln Val Asn His Asp Ile Ala Leu Leu Lys Leu Glu Thr 275 280 285 Pro Val Asn Leu Thr Asp Phe Val Val Pro Ile Cys Leu Pro Glu Lys 290 295 300 Arg Phe Ala Val Tyr Glu Leu Ser Ser Ile Lys Phe Ser Met Val Ser 305 310 315 320 Gly Trp Gly Arg Leu Leu Asp Gly Gly Ala Thr Ser Thr Phe Leu Met 325 330 335 Arg Val His Leu Pro Arg Val Lys Thr Gln Glu Cys Glu Lys Gln Ala 340 345 350 Asn Leu Asn Ile Thr Glu Asn Met Phe Cys Ala Gly Asp Leu Thr Gly 355 360 365 Lys Lys Asp Ser Cys Lys Gly Asp Ser Gly Gly Pro His Ala Thr Lys 370 380 Tyr Lys Asn Thr Trp Phe Leu Thr Gly Ile Val Ser Trp Gly Lys Gly 385 390 395 400 Cys Ala Val Glu Gly Ser Tyr Gly Val Tyr Thr Arg Val Ser Arg Tyr 405 410 415 Ile Asn Trp Leu Lys Arg His Met Glu 420 425 <210> 15 <211> 443 <212> PRT <213> Oryctolagus cuniculus <400> 15 Met Ala Pro Gln Ala Arg Gly Leu Gly Leu Cys Ser Leu Leu Ala Leu $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Sequence Listing
Gln Ala Ser Leu Ala Ala Val Phe Ile Thr Gln Glu Glu Ala His Ser
20 25 30 Val Leu Arg Arg Gln Arg Arg Ala Asn Ser Phe Leu Glu Glu Leu Arg 35 40 45 Pro Gly Ser Leu Glu Arg Glu Cys Lys Glu Glu Leu Cys Ser Phe Glu 50 60 Glu Ala Arg Glu Val Phe Gln Ser Thr Glu Arg Thr Lys Gln Phe Trp 70 75 80 Ile Thr Tyr Asn Asp Gly Asp Gln Cys Ala Ser Asn Pro Cys Gln Asn 85 90 95 Gly Gly Ser Cys Glu Asp Gln Ile Gln Ser Tyr Ile Cys Phe Cys Leu 100 105 110Ala Asp Phe Glu Gly Arg Asn Cys Glu Lys Asn Lys Asn Asp Gln Leu 115 120 125 Val Gly Ser Gln Arg Ser Cys Arg Cys His Glu Gly Tyr Thr Leu Leu 145 150 155 160 Pro Asn Gly Val Ser Cys Thr Pro Thr Val Asp Tyr Pro Cys Gly Lys
165 170 175 Val Pro Ala Leu Glu Lys Arg Gly Ala Ser Asn Pro Gln Gly Arg Ile 180 185 190 Val Gly Gly Lys Val Cys Pro Lys Gly Glu Cys Pro Trp Gln Ala Ala 195 200 205 Leu Met Asn Gly Ser Thr Leu Leu Cys Gly Gly Ser Leu Leu Asp Thr 210 215 220 His Trp Val Val Ser Ala Ala His Cys Phe Asp Lys Leu Ser Ser Leu 225 230 235 240 Arg Asn Leu Thr Ile Val Leu Gly Glu His Asp Leu Ser Glu His Glu 245 250 255 Gly Asp Glu Gln Val Arg His Val Ala Gln Leu Ile Met Pro Asp Lys 265

Tyr Val Pro Gly Lys Thr Asp His Asp Ile Ala Leu Leu Arg Leu Leu 275 280 285

Gln Pro Ala Ala Leu Thr Asn Asn Val Val Pro Leu Cys Leu Pro Glu 290 295 300

Arg Asn Phe Ser Glu Ser Thr Leu Ala Thr Ile Arg Phe Ser Arg Val 315 320

Ser Gly Trp Gly Gln Leu Leu Tyr Arg Gly Ala Leu Ala Arg Glu Leu 325 330 335

Met Ala Ile Asp Val Pro Arg Leu Met Thr Gln Asp Cys Val Glu Gln 340 345 350

Ser Glu His Asn Pro Gly Ser Pro Glu Val Thr Gly Asn Met Phe Cys 355 360 365

Ala Gly Tyr Leu Asp Gly Ser Lys Asp Ala Cys Lys Gly Asp Ser Gly 370 375 380

Gly Pro His Ala Thr Ser Tyr His Gly Thr Tyr Leu Thr Gly Val Val 385 390 395 400

Ser Trp Gly Glu Gly Cys Ala Arg Val Gly His Val Gly Val Tyr Thr 405 410 415

Arg Val Ser Arg Asp Thr Glu Trp Leu Ser Arg Leu Met Arg Ser Lys 420 425 430

Leu His His Gly Ile Gln Arg His Pro Phe Pro 435

<210> 16

<211> 681

<212> PRT

<213> Mus musculus

<400> 16

Met Val Pro Gln Ala His Gly Leu Leu Leu Cys Phe Leu Leu Gln 1 5 10 15

Leu Gln Gly Pro Leu Gly Thr Ala Val Phe Ile Thr Gln Glu Glu Ala 20 25 30

His Gly Val Leu His Arg Gln Arg Arg Ala Asn Ser Leu Leu Glu Glu 35 40 45

Leu Trp Pro Gly Ser Leu Glu Arg Glu Cys Asn Glu Glu Gln Cys Ser 50 60 Phe Glu Glu Ala Arg Glu Ile Phe Lys Ser Pro Glu Arg Thr Lys Gln 65 70 75 80 Phe Trp Ile Val Tyr Ser Asp Gly Asp Gln Cys Ala Ser Asn Pro Cys 85 90 95 Gln Asn Val Gly Thr Cys Gln Asp His Leu Lys Ser Tyr Val Cys Phe 100 105 110Cys Leu Leu Asp Phe Glu Gly Arg Asn Cys Glu Lys Ser Lys Asn Glu 115 120 125 Gln Leu Ile Cys Ala Asn Glu Asn Gly Asp Cys Asp Gln Tyr Cys Arg 130 135 140 Asp His Val Gly Thr Lys Arg Thr Cys Ser Cys His Glu Asp Tyr Thr 145 150 155 160 Leu Gln Pro Asp Glu Val Ser Cys Lys Pro Lys Val Glu Tyr Pro Cys 165 170 175 Gly Arg Ile Pro Val Val Glu Lys Arg Asn Ser Ser Ser Arg Gln Gly 180 185 190 Arg Ile Val Gly Gly Asn Val Cys Pro Lys Gly Glu Cys Pro Trp Gln
195 200 205 Ala Val Leu Lys Ile Asn Gly Leu Leu Leu Cys Gly Ala Val Leu Leu 210 215 220 Asp Ala Arg Trp Ile Val Thr Ala Ala His Cys Phe Asp Asn Ile Arg 225 230 235 240 Tyr Trp Gly Asn Ile Thr Val Val Met Gly Glu His Asp Phe Ser Glu 245 250 255 Lys Asp Gly Asp Glu Gln Val Arg Arg Val Thr Gln Val Ile Met Pro 260 265 270 Asp Lys Tyr Ile Arg Gly Lys Ile Asn His Asp Ile Ala Leu Leu Arg 275 280 285 Leu His Arg Pro Val Thr Phe Thr Asp Tyr Val Val Pro Leu Cys Leu 290 295 300 Page 23

Pro Glu Lys Ser Phe Ser Glu Asn Thr Leu Ala Arg Ile Arg Phe Ser 305 310 315 320 Arg Val Ser Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala Thr Ala Leu 325 330 335 Glu Leu Met Ser Ile Glu Val Pro Arg Leu Met Thr Gln Asp Cys Leu 340 345 350 Glu His Ala Lys His Ser Ser Asn Thr Pro Lys Ile Thr Glu Asn Met 355 360 Phe Cys Ala Gly Tyr Met Asp Gly Thr Lys Asp Ala Cys Ala Gly Asp 370 380 Ser Gly Gly Pro His Ala Thr His Tyr His Gly Thr Trp Tyr Leu Thr 385 390 395 400 Gly Val Val Ser Trp Gly Glu Gly Cys Ala Ala Ile Gly His Ile Gly 405 410 415 Val Tyr Thr Arg Val Ser Gln Tyr Ile Asp Trp Leu Val Arg His Met 420 425 430 Asp Ser Lys Leu Gln Val Gly Val Phe Arg Leu Pro Leu Leu Gly Ser 435 440 445 Ala Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro 450 455 460 Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys 465 470 475 480 Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val 485 490 495 Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr 500 505 510 Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu 515 520 525 Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His 530 540 Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Page 24

545

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln 565 570 575

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu 580 585 590

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro 595 600 605

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn 610 615 620

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu 625 630 635 640

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val 645 650 655

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln 660 670

Lys Ser Leu Ser Leu Ser Pro Gly Lys 675 680

<210> 17

<211> 407

<212> PRT <213> Bos taurus

<400> 17

Ala Asn Gly Phe Leu Glu Glu Leu Leu Pro Gly Ser Leu Glu Arg Glu 10 15

Cys Arg Glu Glu Leu Cys Ser Phe Glu Glu Ala His Glu Ile Phe Arg 20 25 30

Asn Glu Glu Arg Thr Arg Gln Phe Trp Val Ser Tyr Asn Asp Gly Asp 35 40 45

Gln Cys Ala Ser Ser Pro Cys Gln Asn Gly Gly Ser Cys Glu Asp Gln 50 55 60

Leu Arg Ser Tyr Ile Cys Phe Cys Pro Asp Gly Phe Glu Gly Arg Asn 65 70 75 80

Cys Glu Thr Asp Lys Gln Ser Gln Leu Ile Cys Ala Asn Asp Asn Gly Page 25 Gly Cys Glu Gln Tyr Cys Gly Ala Asp Pro Gly Ala Gly Arg Phe Cys 100 105 110 Trp Cys His Glu Gly Tyr Ala Leu Gln Ala Asp Gly Val Ser Cys Ala 115 120 125 Pro Thr Val Glu Tyr Pro Cys Gly Lys Ile Pro Val Leu Glu Lys Arg 130 135 140 Asn Gly Ser Lys Pro Gln Gly Arg Ile Val Gly Gly His Val Cys Pro 145 150 155 160 Lys Gly Glu Cys Pro Trp Gln Ala Met Leu Lys Leu Asn Gly Ala Leu 165 170 175 Leu Cys Gly Gly Thr Leu Val Gly Pro Ala Trp Val Val Ser Ala Ala 180 185 190 His Cys Phe Glu Arg Leu Arg Ser Arg Gly Asn Leu Thr Ala Val Leu 195 200 205 Gly Glu His Asp Leu Ser Arg Val Glu Gly Pro Glu Gln Glu Arg Arg 210 215 220 Val Ala Gln Ile Ile Val Pro Lys Gln Tyr Val Pro Gly Gln Thr Asp 225 230 235 240 His Asp Val Ala Leu Leu Gln Leu Ala Gln Pro Val Ala Leu Gly Asp 245 250 255 His Val Ala Pro Leu Cys Leu Pro Asp Pro Asp Phe Ala Asp Gln Thr 260 265 270 Leu Ala Phe Val Arg Phe Ser Ala Val Ser Gly Trp Gly Gln Leu Leu 275 280 285 Glu Arg Gly Val Thr Ala Arg Lys Leu Met Val Val Leu Val Pro Arg 290 295 300 Leu Leu Thr Gln Asp Cys Leu Gln Gln Ser Arg Gln Arg Pro Gly Gly 305 310 315 Pro Val Val Thr Asp Asn Met Phe Cys Ala Gly Tyr Ser Asp Gly Ser 325 330 335

Lys Asp Ala Cys Lys Gly Asp Ser Gly Gly Pro His Ala Thr Arg Phe 340 345 350

Arg Gly Thr Trp Phe Leu Thr Gly Val Val Ser Trp Gly Glu Gly Cys 355 360 365

Ala Ala Ala Gly His Phe Gly Ile Tyr Thr Arg Val Ser Arg Tyr Thr 370 380

Ala Trp Leu Arg Gln Leu Met Gly His Pro Pro Ser Arg Gln Gly Phe 385 390 395 400

Phe Gln Val Pro Leu Leu Pro 405